

SEQUENCE LISTING

<110> MultiGene Biotech GmbH

<120> Novel retina-specific human proteins C7orf9, C12orf7, MPP4 and F379

<130> M36888US

<150> 60/253,751

<151> 2000-11-29

<160> 45

<170> PatentIn version 3.1

<210> 1

<211> 2435

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> artificial sequence, Translation start at 209; stop at 2435

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<223> genomic DNA, Exon from 1 to 108

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ttcttcagta tggaataatg 320

<210> 3

<211> 512

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 173 to 352

<400> 3

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tgtagcacag ggcttggtt gctgaggaaa tgctattgaa aatatattcc agtgtgctga 180

gagctggtgg ccagtgggac tgagtgagct gtgtgcctg tattgaccg cttcctagtc 240

ctgaattcct ttcagaagct ccggcaggga ggatgatata gtcagacaaa ggagcagatc 300

caccagacaa gaaggacatg aagctttcta cagccaccaa tccacagaat ggtatgtgtc 360

accaggactc cttttctaga ccagaaagta atatcacctc tgacatgtga tcaaataaat 420

aggcagaaat cctgacagac ttactgtgat ccctatgagg atcttgtaca tttttggttg 480

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<210> 4

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 165 to 286

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acacccgcac ctgccacccc atctctgctc tcttcctttc ctaggcctct cccagatcct 180

gaggcttgat ctgcaagagc tgagtctggt ctacagcaga gatgtgaatg gagtgtgtct 240

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gctcggaagc ctttgcttgc tgaagggggt gtggggagtg ttagaaaaat gacagcttca 360

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ctgtaggccca ggaagtccaa aatccagt

448

<210> 5

<211> 448

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 206 to 283

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cacaacttac taaacaccaa ccacaccgtg ctgtgcagcc attggtgcag ttgcctgggg 120

tgtttcttct ctttgagagt cttaaatcca aaatggcaat agtcatatta tcaatatcaa 180

ttctccctcc cttgtccttc tgcagattta tgactgcctc caggaattta aagaaaagaa 240

actagttcct gccacaccac atgcacaggt gttatcctat gaggtaagga gattttattc 300

cacaggatag tagagctctg atgtggtgcc attttcccca cattgctagt tcaaatgaat 360

taaaggttct aaggaaaagt tttattgatg actatgcac taataaatgt ttctaattga 420

actttaatat aaggaagaac attggctg 448

<210> 6

<211> 384

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 165 to 245

<400> 6

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agatttttta aaatttttat aatgtatcct tttccatgaa ccaggtagtg gagttattac 180

gtgaaacccc tacttcccct gagatccaag agctgagaca aatgctccag gctccacact 240

tcaaggcaag tgcttgctaa aatagaaaag atgtcccat ctggcacata gacaaagttg 300

ggaaggagaa atatatgtga tggaaaatgt tctctctgaa tagatgttct attactgtac 360

acggttactg accaacagat tgta 384

<210> 7

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 133 to 264

<400> 7

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ccctgtgcc aggccttgct cagtgccat gacacgatag ctgagaaaga ttttgaacco 180
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agcttaggtt aattgtgaac caaattatat ctagtggtta cttgggcagt agccttgcc 360
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aaacatatta catgtcacgg tgttttct 448

<210> 8

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 166 to 247

<400> 8

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ccagagctgc attgattgaa agaccagagc tgcattgatt gaggggaagcc acctggaaaa 120

tggatcatgtc aggtaacaga gggatctcgt ctattctctc ttcagggagc caccatcaag 180

cgccacgaga tgacagggga catcttggtg gccaggatca tccacggtgg gctggcggag 240

agaagtggta agctggagca gctgggattg agagttacca gaaaaacagg aaacccttga 300

ctgttttaggc ttctttctag agaaatccct tttttttctt tttttttttc tttttttttt 360

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<210> 9

<211> 448

<212> DNA

<213> Homo sapiens

<220>

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<223> genomic DNA, Exon from 162 to 247

<400> 9

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ttctttgtat ctttccttgt tttttgttac tgtctgccta gggttgctat atgctggaga 180

caaaactgga gaagtgaatg gagtttcagt tgagggactg gaccctgaac aagtgatcca 240

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actgggacca tcaagccac gtgtgtgcac tgggatgtac cggggactca agttctcttg 360

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<210> 10

<211> 384

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 158 to 229

<400> 10

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ccatgattgt catcctttct cctctcttat ttccaggcc atgtctcgag gcacaatcat 180

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ctgagccttc aatctcacac acagtaaadc cccaagtaac agcaactaaa tatgatgcgt 300

aataatccta tcctttgtac tgtgttggac ctggattcaa gactgtgttg gatatttttc 360

aatactgatg gcccgagaag caaa 384

<210> 11

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 138 to 334

<400> 11

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<210> 12

<211> 320

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 152 to 216

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 gaggtctggcc atgccacatg 320

<210> 13

<211> 320

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 161 to 178

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 aagaaatagt atttaggaaa aaactcttat ctccaaagtc ttttagaaat ttcttgtagt 240
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 gtgtgatagt cattattaag 320

<210> 14

<211> 384

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 179 to 217

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<223> genomic DNA, Exon from 110 to 130

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ttttgatgca atttatacac atatttataa taactgttta aatatatcaa cattaaaaaa 240

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tttattttta attttaattt 320

<210> 16

<211> 320

<212> DNA

<213> Homo sapiens

<220>

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<221> misc_feature

<223> genomic DNA, Exon from 174 to 188

<400> 16

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tcttaaaggc ttaacttttt catgtctggt tctgcactta cccaaatatt cagaggaact 180

ttcagaaggt aattgttttt atttcctaga tataccaaat agaactatgt ttaagattct 240

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ttgttggtta tggttttccc 320

<210> 17

<211> 320

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 170 to 211

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<211> 512

<212> DNA

<213> Homo sapiens

<400> 18

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512

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<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 160 to 240

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cacatctgat gatttctgtg tgtgactttt tgtgtttagg accctctggg gttggagtaa 180

atgagctcag aagacaactt attgaattta atcccagcca ttttcaaagt gctgtgccac 240

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agtatttcta accatattta tttt 384

<210> 20

<211> 448

<212> DNA

<213> Homo sapiens

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<221> misc_feature

<223> genomic DNA, Exon from 200 to 293

<400> 20

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agatatcttt attcaaatgc atattggtaa tcaaagaatt ctgaagacac tgaaaccttt 180

cattcccttt ttctgataga cactactcgt actaaaaaga gttacgaaat gaatgggcgt 240

gagtatcact atgtgtccaa ggaaacattt gaaaacctca tatatagtca caggtaaagt 300

agagggtcag aagctgattc ttacctcttg ttgttttaca tttgaaatag attccctatt 360

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atgtacatcg atatacagca caccaact 448

<210> 21

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 133 to 241

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<210> 22

<211> 448

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 164 to 298

<400> 22

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aaaacaaaca aaaaataaat gtgcatttaa attttctgtg taggatattc aaggggttcg 180

aacctatgaa ctgaagccct atgtcatatt tataaagcca tcgaatatga ggtgtatgaa 240

acaatctcgg aaaaatgcc aaggtattac tgactactat gtggacatga agttcaaggt 300

aagagcaagt caaaaactac tgtattgctt tcagtggcct ctgcgtggga gagatctggg 360

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caaaaggccc tcaataaaat ggtttact 448

<210> 23

<211> 704

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 197 to 704

<400> 23

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acagcataac aaactgtatt ttttccatth gtccaattaa gtctgtacta tccatatttt 180

tctattttct ctaaaggatg aagacctaca agagatggaa aatttagccc aaagaatgga 240

aactcagttt ggccaatttt ttgatcatgt gattgtgaat gacagcttgc acgatgcatg 300

tgcccagttg ttgtctgcc aacagaaggc tcaggaggag cctcagtggg taccagcaac 360

atggatttcc tcagatactg agtctcaatg agacttcttg tttaatgctg gagttttaac 420

actgtaccct tgatacagcg atccatagtt gcaatctaaa acaacagtat ttgaccatt 480

ttaatgtgta caactttaaa agtgcagcaa tttattaatt aatcttattt gaaaaaaatt 540

tttattgtat ggttatgtgg ttacctattt taacttaatt ttttttcctt tacctcatat 600

gcagctgtgg tagaaatatg aataatgtta agtcactgag tatgagaacc tttcgcatat 660

ttcacatgat cttttttaaga tttaaataaa gagctttcct aaat 704

<210> 24

<211> 637

<212> PRT

<213> Homo sapiens

<400> 24

Met Ile Gln Ser Asp Lys Gly Ala Asp Pro Pro Asp Lys Lys Asp Met
1 5 10 15

Lys Leu Ser Thr Ala Thr Asn Pro Gln Asn Gly Leu Ser Gln Ile Leu
20 25 30

Arg Leu Val Leu Gln Glu Leu Ser Leu Phe Tyr Ser Arg Asp Val Asn
35 40 45

Gly Val Cys Leu Leu Tyr Asp Leu Leu His Ser Pro Trp Leu Gln Ala
50 55 60

Leu Leu Lys Ile Tyr Asp Cys Leu Gln Glu Phe Lys Glu Lys Lys Leu
65 70 75 80

Val Pro Ala Thr Pro His Ala Gln Val Leu Ser Tyr Glu Val Val Glu
85 90 95

Leu Leu Arg Glu Thr Pro Thr Ser Pro Glu Ile Gln Glu Leu Arg Gln
100 105 110

Met Leu Gln Ala Pro His Phe Lys Ala Leu Leu Ser Ala His Asp Thr
115 120 125

Ile Ala Gln Lys Asp Phe Glu Pro Leu Leu Pro Pro Leu Pro Asp Asn
130 135 140

Ile Pro Glu Ser Glu Glu Ala Met Arg Ile Val Cys Leu Val Lys Asn
145 150 155 160

Gln Gln Pro Leu Gly Ala Thr Ile Lys Arg His Glu Met Thr Gly Asp
165 170 175

Ile Leu Val Ala Arg Ile Ile His Gly Gly Leu Ala Glu Arg Ser Gly
180 185 190

Leu Leu Tyr Ala Gly Asp Lys Leu Val Glu Val Asn Gly Val Ser Val
195 200 205

Glu Gly Leu Asp Pro Glu Gln Val Ile His Ile Leu Ala Met Ser Arg
210 215 220

Gly Thr Ile Met Phe Lys Val Val Pro Val Ser Asp Pro Pro Val Asn
225 230 235 240

Ser Gln Gln Met Val Tyr Val Arg Ala Met Thr Glu Tyr Trp Pro Gln
245 250 255

Glu Asp Pro Asp Ile Pro Cys Met Asp Ala Gly Leu Pro Phe Gln Lys
260 265 270

Gly Asp Ile Leu Gln Ile Val Asp Gln Asn Asp Ala Leu Trp Trp Gln
275 280 285

Ala Arg Lys Ile Ser Asp Pro Ala Thr Cys Ala Gly Leu Val Pro Ser
290 295 300

Asn His Leu Leu Lys Arg Lys Gln Arg Glu Phe Trp Trp Ser Gln Pro
305 310 315 320

Tyr Gln Pro His Thr Cys Leu Lys Ser Thr Leu Ser Ile Ser Met Glu
325 330 335

Glu Glu Asp Asp Met Lys Ile Asp Glu Lys Cys Val Glu Ala Asp Glu
340 345 350

Glu Thr Phe Glu Ser Glu Glu Leu Ser Glu Asp Lys Glu Glu Phe Val
355 360 365

Gly Tyr Gly Gln Lys Phe Phe Ile Ala Gly Phe Arg Arg Ser Met Arg
370 375 380

Leu Cys Arg Arg Lys Ser His Leu Ser Pro Leu His Ala Ser Val Cys
385 390 395 400

Cys Thr Gly Ser Cys Tyr Ser Ala Val Gly Ala Pro Tyr Glu Glu Val
405 410 415

Val Arg Tyr Gln Arg Arg Pro Ser Asp Lys Tyr Arg Leu Ile Val Leu
420 425 430

Met Gly Pro Ser Gly Val Gly Val Asn Glu Leu Arg Arg Gln Leu Ile
435 440 445

Glu Phe Asn Pro Ser His Phe Gln Ser Ala Val Pro His Thr Thr Arg
450 455 460

Thr Lys Lys Ser Tyr Glu Met Asn Gly Arg Glu Tyr His Tyr Val Ser
465 470 475 480

Lys Glu Thr Phe Glu Asn Leu Ile Tyr Ser His Arg Met Leu Glu Tyr
485 490 495

Gly Glu Tyr Lys Gly His Leu Tyr Gly Thr Ser Val Asp Ala Val Gln
500 505 510

Thr Val Leu Val Glu Gly Lys Ile Cys Val Met Asp Leu Glu Pro Gln
515 520 525

Asp Ile Gln Gly Val Arg Thr His Glu Leu Lys Pro Tyr Val Ile Phe
530 535 540

Ile Lys Pro Ser Asn Met Arg Cys Met Lys Gln Ser Arg Lys Asn Ala
545 550 555 560

Lys Val Ile Thr Asp Tyr Tyr Val Asp Met Lys Phe Lys Asp Glu Asp
565 570 575

Leu Gln Glu Met Glu Asn Leu Ala Gln Arg Met Glu Thr Gln Phe Gly
580 585 590

Gln Phe Phe Asp His Val Ile Val Asn Asp Ser Leu His Asp Ala Cys
595 600 605

Ala Gln Leu Leu Ser Ala Ile Gln Lys Ala Gln Glu Glu Pro Gln Trp
610 615 620

Val Pro Ala Thr Trp Ile Ser Ser Asp Thr Glu Ser Gln
625 630 635

<210> 25

<211> 1190

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> artificial sequence, Translation start at 48, stop at 638

ataaacattg ggctgcacat agagacttaa ttttagattt agacaaaatg gaaattattt 60

catcaaaaact attcatttta ttgacttttag ccacttcaag cttgttaaca tcaaacattt 120

tttgtgcaga tgaattagtg atstccaatc ttcacagcaa agaaaattat gacaaatatt 180

ctgagcctag aggataccca aaaggggaaa gaagcctcaa ttttgaggaa ttaaagatt 240

ggggacccaaa aaatgttatt aagatgagta cacctgcagt caataaaatg ccacactcct 300

tgcccaactt gccattgaga tttgggagga acgttcaaga agaaagaagt gctggagcaa 360

cagccaacct gcctctgaga tctggaagaa atatggaggt gagcctcgtg agacgtgttc 420

ctaacctgcc ccaaaggttt gggagaacaa caacagccaa aagtgtctgc aggatgctga 480

gtgatttgtg tcaaggatcc atgcattcac catgtgccaa tgacttattt tactccatga 540

cctgccagca ccaagaaatc cagaatcccg atcaaaaaca gtcaaggaga ctgctattca 600

agaaaataga tgatgcagaa ttgaaacaag aaaaataaga aacctggagc ctgtccctaa 660

agctgtggcc tgtaatctac aaatggctct atagcgaaga ccacacggaa gagtagctac 720

atacacttca tcagctatgg atcatcaacg gcaatttttc cttgtcagta cagctataat 780

agtatcttga aagttgtaaa aaaattaaag catatttggt acgtaaagtt aaaatgattt 840

ttgtctgaat aaaaaaaaaag cattgcaaat gctttagaaa tctctgataa tggagagaga 900

gacagaggac cctcctcact accctatata aaaatcattg gcacagttac acttaataaa 960

aaaaattaaa cagaagagca ccttgaaaaa cattatgatg gaaattaaat agtatgccag 1020

aataacatgg ttgacaaata agtgaacaag gattaataat cacttacaaa cgtgtttctg 1080

tacacccttt ctatcgtgtc aaatgttaat gaatctgtga tcaattgaaa tgtaaattgtc 1140

tgtgtaaaac tacaaaataa aaactcttag acttttagga gaaaagaaaa 1190

<210> 26

<211> 256

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 1 to 185

<400> 26

ataaacattg ggctgcacat agagacttaa ttttagattt agacaaaatg gaaattattt 60

catcaaaact attcatttta ttgacttttag ccacttcaag cttgttaaca tcaaacattt 120

tttgtgcaga tgaattagtg atstccaatc ttcacagcaa agaaaattat gacaaatatt 180

ctgaggtaag ttttttaa atctcttaatg tgagtagcat taattacata atattaatcc 240

taagtcta at gatttt 256

<210> 27

<211> 512

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 62 to 462

<400> 27

gggttttaa at ctgttgctta taacaacagt atgttattgt aatggtcatt tctaattata 60

gcctagagga tacccaaaag gggaaagaag cctcaatttt gaggaattaa aagattgggg 120

accaaaaaat gttattaaga tgagtacacc tgcagtcaat aaaatgccac actccttcgc 180

caacttgcca ttgagatttg ggaggaacgt tcaagaagaa agaagtgctg gagcaacagc 240

caacctgcct ctgagatctg gaagaaatat ggaggtgagc ctctgagagc gtgttcctaa 300

cctgccccaa aggtttggga gaacaacaac agccaaaagt gtctgcagga tgctgagtga 360

tttgtgtcaa ggatccatgc attcaccatg tgccaatgac ttattttact ccatgacctg 420

ccagcaccaa gaaatccaga atcccgatca aaaacagtca aggtaaatac ctggaaacca 480

gtcaaagtgc atgggcagtt atatagaggt gg 512

<210> 28

<211> 768

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 115 to 718

<400> 28

acacaattca actcaagtat aattaggcag ttaggactat ggcttgtatt tgtatacaca 60

cttgcattgct gttgttctga tgggtgacaa cattttatac tgcttacatt ttaggagact 120

gctattcaag aaaatagatg atgcagaatt gaaacaagaa aaataagaaa cctggagcct 180

gtccctaaag ctgtggcctg taatctacaa atggctctat agcgaagacc acacggaaga 240

gtagctacat acacttcacg agctatggat catcaacggc aatttttctt tgtcagtaca 300

gctataatag tatcttgaaa gttgtaaaaa aattaaagca tatttgttac gtaaagttaa 360

aatgattttt gtctgaataa aaaaaaagca ttgcaaatgc ttagaaaatc tctgataatg 420

gagagagaga cagaggaccc tcctcactac cctatataaa aatcattggc acagttacac 480

ttaataaaaa aaattaaaca gaagagcacc ctgaaaaaca ttatgatgga aattaaatag 540

tatgccagaa taacatgggt gacaaataag tgaacaagga ttaaaaatca cttacaaacg 600

tgtttctgta caccctttct atcgtgtcaa atgttaatga atctgtgatc aattgaaatg 660

taaatgtctg tgtaaaaacta caaaataaaa actcttagac tttagggaga aaagaaaaag 720

gcaactatga gttacctctt ttagtgtctc ctctatctac atccagaa

768

<210> 29

<211> 196

<212> PRT

<213> Homo sapiens

<400> 29

Met Glu Ile Ile Ser Ser Lys Leu Phe Ile Leu Leu Thr Leu Ala Thr
1 5 10 15

Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Ala Asp Glu Leu Val Ile
20 25 30

Ser Asn Leu His Ser Lys Glu Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
35 40 45

Gly Tyr Pro Lys Gly Glu Arg Ser Leu Asn Phe Glu Glu Leu Lys Asp
50 55 60

Trp Gly Pro Lys Asn Val Ile Lys Met Ser Thr Pro Ala Val Asn Lys
65 70 75 80

Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Val
85 90 95

Gln Glu Glu Arg Ser Ala Gly Ala Thr Ala Asn Leu Pro Leu Arg Ser
100 105 110

Gly Arg Asn Met Glu Val Ser Leu Val Arg Arg Val Pro Asn Leu Pro
115 120 125

Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Val Cys Arg Met Leu
130 135 140

Ser Asp Leu Cys Gln Gly Ser Met His Ser Pro Cys Ala Asn Asp Leu
145 150 155 160

Phe Tyr Ser Met Thr Cys Gln His Gln Glu Ile Gln Asn Pro Asp Gln
165 170 175

Lys Gln Ser Arg Arg Leu Leu Phe Lys Lys Ile Asp Asp Ala Glu Leu
180 185 190

Lys Gln Glu Lys
195

<210> 30

<211> 1188

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> artificial sequence, Translation start at 347, stop at 604

<400> 30

acacacaacg gggtttcggg gctgtggacc ctgtgccagg aaaggaaggg cgcagctcct 60

gcaatgcgga gcagccaggg cagtgggcac caggcttttag cctccctttc tcaccctaca 120

gagggcaggc ctttcagctc cattctcctc caaggctgca gagggggcag gaattggggg 180

tgacaggaga gctgtaaggt ctccagtggg tcattctggg ccagagatg ggtgctgaag 240

ctcccacgcc tgcctgtgaa aatggagtcc tctctcacct gggagagcca ggtgctgccc 300

cgagaaggat gcatttatgg cttcttgaag tctttcctga cccccgatgc tgctgactat 360

agagacaaag tctcactatg ttgctcaggc tggctctgaa ctctggcct caagcgatcc 420

tcccacctya gcctcccaa gwgttgggat tatagacatg agccactgca cctggccgac 480

cttgggcaag ttcttaaacc cttcaaagcc tcatttttct ccaatcayaa aagggaaaga 540

tggtaatatt ttccccwcca aattcttgtc ggatgccctc acagaattga gattatgtac 600

gtaaaacacc aggtgcctaa cccggcacag agcaggaggg ctaagcgtga catccagcac 660

gtggtcagtg gaatccagta ttcctacca cctctctagt ctcccccca cccctctccc 720

tttcagaggc accaagctgc ttgtggtett gtctattccc actccctgcc tgactgaaca 780

ttttctccac ctctgatca tcagcagcag aaactggctg ctcttcctcc tgggtagaca 840

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gccagactgt atttcccagc tgcccctgca gtgagatgtg gccatcggag ccagcattgg      900

ccaatggact ctgcatggga gtgacgcatg cwgcctccag gcttgtccct aaaacctccc      960

acgtgtcctc sgctgtctct tcccacytcc aaggagcacg gcaattgtgg aagaccaga      1020

ttagtgatgg cagaaccata gatggggagga acctgggtcc ctgacttaaa gtatcatgga      1080

tttgatggtt cccttagtga gaaataaaact tccattgtgt ttaagccttt atttgtttat      1140

agttggttac agcaactgcc ttcttttaat taaaacactc ctgctgct                      1188

```

<210> 31

<211> 85

<212> PRT

<213> Homo sapiens

<400> 31

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Met Leu Leu Thr Ile Glu Thr Lys Ser His Tyr Val Ala Gln Ala Gly
1           5           10           15

```

```

Leu Glu Leu Leu Ala Ser Ser Asp Pro Pro Thr Ser Ala Ser Gln Ser
          20           25           30

```

```

Val Gly Ile Ile Asp Met Ser His Cys Thr Trp Pro Thr Leu Gly Lys
          35           40           45

```

Phe Leu Asn Pro Ser Lys Pro His Phe Ser Pro Ile Thr Lys Gly Lys
50 55 60

Asp Gly Asn Ile Phe Pro Thr Lys Phe Leu Ser Asp Ala Leu Thr Glu
65 70 75 80

Leu Arg Leu Cys Thr
85

<210> 32

<211> 560

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 101 to 460

<400> 32

tatatgggaa tgagccagct gcaccgctgc tgacagtggc tgggataatc ctccctgagc 60

tgttccaagg attagtcctg ctgccctgtg cccagctccc acacaacggg gtttcggggc 120

tgtggaccct gtgccaggaa aggaagggcg cagctcctgc aatgcggagc agccagggca 180

gtgggcacca ggcttttagcc tccctttctc accctacaga gggcaggccc ttcagctcca 240

ttctcctcca aggctgcaga gggggcagga attgggggtg acaggagagc tgtaaggctct	300
cagtggggtc attctgggcc cagagatggg tgctgaagct cccacgcctg cctgtgaaaa	360
tggagtcttc tctcacctgg gagagccagg tgctgccccg agaaggatgc atttatggct	420
tcatgaagtc tttcctgacc ccgatgctg ctgactatag gtaagtctga gcaaactctgg	480
gggagcctca tcttggcatg agaaagagat ggcttcttct aagccactg gccgtgatcc	540
caggattata acacattctg	560

<210> 33

<211> 405

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 101 to 305

<400> 33

catgagaggt agtataatat agaggatatg tgtgcttact aagaggctgc ctgtctgacc	60
ttggacaagt tctttttatt tatttattta ttttttatag agacaaagtc tcaactatgtt	120
gtcaggctg gtcttgaact cctggcctca agcgatcctc ccaccttagc ctcccaaaga	180
gttgggatta tagacatgag cactgcacc tggccgacct tgggcaagtt cttaaaccct	240

tcaaagcctc atttttctcc aatcataaaa gggaaagatg gtaatatattt cccctccaaa 300

ttcttgtaag tattaacat tgtatatgta ttttgaacac gattaagctc taaacacttg 360

ttaggaagca ggagtagcat ttgaaacaaa cagctctttt cccac 405

<210> 34

<211> 821

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 101 to 721

<400> 34

aagtattaaa cattgtatat gtattttgaa cagattaag ctctaaacac ttgttaggaa 60

gcaggagtag catttgaaac aaacagctct tttccacag gtcggatgcc ctcacagaat 120

tgagattatg tacgtaaaac accaggtgcc taaccggca cagagcagga gggctaagcg 180

tgacatccag cacgtggtca gtggaatcca gtattcctac ccacctctct agtctccct 240

ccacctctct cctttcaga ggcaccaagc tgcttggtgt cttgtctatt cccactccct 300

gctgactga acattttctc cacctcctga tcatcagcag cagaaactgg ctgctcttcc 360

tcctgggtag acagccagac tgtatttccc agctgccct gcagtgagat gtggccatcg	420
gagccagcat tggccaatgg actctgcatg ggagtgaagc atgctgcctc caggcttgtc	480
cctaaaacct cccacgtgtc ctccgctgc tcttcccact tccaaggagc acggcaattg	540
tggaagaccc agattagtga tggcagaacc atagatggga ggaacctggg tccctgactt	600
aaagtatcat ggatttggat gttcccttag tgagaaataa acttccattg tgtttaagcc	660
tttatttgtt tatagttggg tacagcaact gccttctttt aattaaaaca ctctgctgc	720
ttcatgttgc tggaatgctt gtaaccctgc cctgcttcac cagggttaact cctacttggc	780
ctttaagttt atctctgctg tcacaccgtc cagaaagcct t	821

<210> 35

<211> 1514

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> artificial sequence, Translation start at 155, stop at 1192

<400> 35

gaaagtccag ccattctgtta cctgcgttgc ttcttggggr gggatagtcc acctggaggc	60
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attcggagac ccagtgattg tgctccgygg agcctggggt gtgccccgcg ttgactgcct	120
---	-----

catagatacc ctacgaacc caaatgccag ctgcatgaga aaagggactc accttctggt	180
tccctgctg gaagaggaag agctggcatt gcacaggaga cggctggaca tgtctgaggc	240
actgccctgc ccgggcaagg agacccccac cccaggctgc aggctggggg ccctgtattg	300
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cggttccag agtggttggt ccctgctcag ccactgtcct ttccttgatg tgaaccagca	480
ggacaaagga ggggacacgg ccctcatggt ggtgcccac gcaggccacg tgctctagt	540
gagtctctg ctcaactact atgtgggcct ggacctggaa cgccgggacc agcgggggct	600
cacggcggtta atgaaggctg ccatgcggaa ccgctgtgct gacctgacag cagtggacct	660
tgttcggggc aagacggccc tggaatgggc agtgctgacc gacagcttcg acaccgtgtg	720
gaggattcgg cagctgctga ggcgcccca agtgaggcag cttagccagc actacaagcc	780
cgagtggccg gccttgctcg ggctcgtggc ccaggcccag gccaggccc aggttgcccc	840
ttcactccta gaacggctgc aggctacctt gagectccc tttgcccggt ctctcagga	900
ggggggtggt ctggaccacc ttgtgactgc cacaaccage ctggccagtc ccttcgtcac	960
cactgcctgc cacactctgt gcctgacca tccaccttcg ctgggcacc gaagcaagtc	1020
cgtgccagag ctgttagtgc cagccgaagc ccagtccttc aggacaccaa agtctggccc	1080
ttcctctctg gcgataccag gagctcagga tagagaagag gaaacaggag gaggaggcca	1140
gaatggcaca gaagtagggg aagatgggat aggacaggct gggaacaggt aatcaggccc	1200

ctcccagggc ttctttcccc tctggagtgc ctccggcctc cccatccacc tctgcctaag 1260

taaatctgct ctcaacctat atatatacaa ggtcattcat tctagcattg ttgcaagag 1320

tgaaagagtg gaaacacccg aagtgtccat cagtaaggga caggctagat tgattacgga 1380

tgtaattgct gtccatccat acagagcata ctctacagtg tattctaaaa taagactaag 1440

gaagctgttt atattctgat atgaaactac catcaagatg tataaagtaa aaataactaa 1500

ggagtgggaac agtg 1514

<210> 36

<211> 1544

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> artificial sequence, Translation start at 155, stop at 1222

<400> 36

gaaagtccag ccatctgtta cctgcgttgc ttctctggggr gggatagtcc acctggaggc 60

attcggagac ccagtgattg tgctccgygg agcctgggct gtgccccgcy ttgactgcct 120

catagatacc ctacgaaccc caaatgccag ctgcatgaga aaagggactc accttctggt 180

tccctgcctg gaagaggaag agctggcatt gcacaggaga cggctggaca tgtctgaggc	240
actgccctgc cggggcaagg agacccccac ccaggctgc aggtggggg ccctgtattg	300
ggcctgtgtc cacaatgac ccaccagct ccaagccata ctggatggtg gggctctccc	360
agaggaggcc acccaggtgg acagcaatgg gaggacaggc ctcattggtg catgctacca	420
cggcttccag agtggtgtgg ccctgtcag ccactgtcct ttcttgatg tgaaccagca	480
ggacaaagga ggggacacgg ccctcatgtt ggctgcccc gcaggccacg tgctctagt	540
gagtctcctg ctcaactact atgtgggcct ggacctggaa cgccgggacc agcgggggct	600
cacggcgta atgaaggctg ccattgcggaa ccgtgtgag tgcgtggcca cctcctcat	660
ggcaggtgct gacctgacag cagtggaccc tgttcggggc aagacggccc tggaatggg	720
agtgtgacc gacagcttcg acaccgtgtg gaggattcgg cagtgtctga ggcggcccca	780
agtggagcag cttagccagc actacaagcc cgagtggccg gccttgctcg ggctcgtggc	840
ccaggcccag gcccaggccc aggttgcccc ttcaactcta gaacggctgc aggtacctt	900
gagcctcccc ttgccccgt ctctcagga ggggggtgtt ctggaccacc ttgtgactgc	960
cacaaccagc ctggccagtc cttcgtcac cactgcctgc cacactctgt gccctgacca	1020
tccaccttcg ctgggcaccc gaagcaagtc cgtgccagag ctgttagtgc cagccgaagc	1080
ccagtccttc aggacaccaa agtctggccc ttctctctg gcgataccag gagctcagga	1140
tagagaagag gaaacaggag gaggaggcca gaatggcaca gaagtagggg aagatgggat	1200
aggacaggct gggaacaggc aatcaggccc ctcccagggc ttctttcccc tctggagtgc	1260

ctccggcctc cccatccacc tctgcctaag taaatctgct ctcaacctat atatatacaa 1320

ggtcattcat tctagcattg tttgcaagag tgaaagagtg gaaacacccg aagtgtccat 1380

cagtaaggga caggctagat tgattacgga tgtaattgct gtccatccat acagagcata 1440

ctctacagtg tattctaaaa taagactaag gaagctgttt atattctgat atgaaactac 1500

catcaagatg tataaagtaa aaataactaa ggagtggaac agtg 1544

<210> 37

<211> 345

<212> PRT

<213> Homo sapiens

<400> 37

Met Arg Lys Gly Thr His Leu Leu Val Pro Cys Leu Glu Glu Glu Glu
1 5 10 15

Leu Ala Leu His Arg Arg Arg Leu Asp Met Ser Glu Ala Leu Pro Cys
20 25 30

Pro Gly Lys Glu Thr Pro Thr Pro Gly Cys Arg Leu Gly Ala Leu Tyr
35 40 45

Trp Ala Cys Val His Asn Asp Pro Thr Gln Leu Gln Ala Ile Leu Asp
50 55 60

Gly Gly Val Ser Pro Glu Glu Ala Thr Gln Val Asp Ser Asn Gly Arg
65 70 75 80

Thr Gly Leu Met Val Ala Cys Tyr His Gly Phe Gln Ser Val Val Ala
85 90 95

Leu Leu Ser His Cys Pro Phe Leu Asp Val Asn Gln Gln Asp Lys Gly
100 105 110

Gly Asp Thr Ala Leu Met Leu Ala Ala Gln Ala Gly His Val Pro Leu
115 120 125

Val Ser Leu Leu Leu Asn Tyr Tyr Val Gly Leu Asp Leu Glu Arg Arg
130 135 140

Asp Gln Arg Gly Leu Thr Ala Leu Met Lys Ala Ala Met Arg Asn Arg
145 150 155 160

Cys Ala Asp Leu Thr Ala Val Asp Pro Val Arg Gly Lys Thr Ala Leu
165 170 175

Glu Trp Ala Val Leu Thr Asp Ser Phe Asp Thr Val Trp Arg Ile Arg
180 185 190

Gln Leu Leu Arg Arg Pro Gln Val Glu Gln Leu Ser Gln His Tyr Lys
195 200 205

Pro Glu Trp Pro Ala Leu Ser Gly Leu Val Ala Gln Ala Gln Ala Gln
210 215 220

Ala Gln Val Ala Pro Ser Leu Leu Glu Arg Leu Gln Ala Thr Leu Ser
225 230 235 240

Leu Pro Phe Ala Pro Ser Pro Gln Glu Gly Gly Val Leu Asp His Leu
245 250 255

Val Thr Ala Thr Thr Ser Leu Ala Ser Pro Phe Val Thr Thr Ala Cys
260 265 270

His Thr Leu Cys Pro Asp His Pro Pro Ser Leu Gly Thr Arg Ser Lys
275 280 285

Ser Val Pro Glu Leu Leu Val Pro Ala Glu Ala Gln Ser Phe Arg Thr
290 295 300

Pro Lys Ser Gly Pro Ser Ser Leu Ala Ile Pro Gly Ala Gln Asp Arg
305 310 315 320

Glu Glu Glu Thr Gly Gly Gly Gly Gln Asn Gly Thr Glu Val Gly Glu
325 330 335

Asp Gly Ile Gly Gln Ala Gly Asn Arg
340 345

<210> 38

<211> 355

<212> PRT

<213> Homo sapiens

<400> 38

Met Arg Lys Gly Thr His Leu Leu Val Pro Cys Leu Glu Glu Glu Glu

1 5 10 15

Leu Ala Leu His Arg Arg Arg Leu Asp Met Ser Glu Ala Leu Pro Cys

20 25 30

Pro Gly Lys Glu Thr Pro Thr Pro Gly Cys Arg Leu Gly Ala Leu Tyr

35 40 45

Trp Ala Cys Val His Asn Asp Pro Thr Gln Leu Gln Ala Ile Leu Asp

50 55 60

Gly Gly Val Ser Pro Glu Glu Ala Thr Gln Val Asp Ser Asn Gly Arg

65 70 75 80

Thr Gly Leu Met Val Ala Cys Tyr His Gly Phe Gln Ser Val Val Ala

85 90 95

Leu Leu Ser His Cys Pro Phe Leu Asp Val Asn Gln Gln Asp Lys Gly

100

105

110

Gly Asp Thr Ala Leu Met Leu Ala Ala Gln Ala Gly His Val Pro Leu
115 120 125

Val Ser Leu Leu Leu Asn Tyr Tyr Val Gly Leu Asp Leu Glu Arg Arg
130 135 140

Asp Gln Arg Gly Leu Thr Ala Leu Met Lys Ala Ala Met Arg Asn Arg
145 150 155 160

Cys Glu Cys Val Ala Thr Leu Leu Met Ala Gly Ala Asp Leu Thr Ala
165 170 175

Val Asp Pro Val Arg Gly Lys Thr Ala Leu Glu Trp Ala Val Leu Thr
180 185 190

Asp Ser Phe Asp Thr Val Trp Arg Ile Arg Gln Leu Leu Arg Arg Pro
195 200 205

Gln Val Glu Gln Leu Ser Gln His Tyr Lys Pro Glu Trp Pro Ala Leu
210 215 220

Ser Gly Leu Val Ala Gln Ala Gln Ala Gln Ala Gln Val Ala Pro Ser
225 230 235 240

Leu Leu Glu Arg Leu Gln Ala Thr Leu Ser Leu Pro Phe Ala Pro Ser

245

250

255

Pro Gln Glu Gly Gly Val Leu Asp His Leu Val Thr Ala Thr Thr Ser

260

265

270

Leu Ala Ser Pro Phe Val Thr Thr Ala Cys His Thr Leu Cys Pro Asp

275

280

285

His Pro Pro Ser Leu Gly Thr Arg Ser Lys Ser Val Pro Glu Leu Leu

290

295

300

Val Pro Ala Glu Ala Gln Ser Phe Arg Thr Pro Lys Ser Gly Pro Ser

305

310

315

320

Ser Leu Ala Ile Pro Gly Ala Gln Asp Arg Glu Glu Glu Thr Gly Gly

325

330

335

Gly Gly Gln Asn Gly Thr Glu Val Gly Glu Asp Gly Ile Gly Gln Ala

340

345

350

Gly Asn Arg

355

<210> 39

<211> 183

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 1 to 143

<400> 39

gaaagtccag ccatctgtta cctgcgttgc ttctctggggr gggatagtcc acctggagggc 60

attcggagac ccagtgattg tgctccgygg agcctgggct gtgccccgcg ttgactgcct 120

catagatacc ctacgaaccc caagtaagaa aaaacgacga ccctctctcc gtgagtctca 180

ctg 183

<210> 40

<211> 462

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 108 to 358

<400> 40

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taggatccag ggagacactc actactcctc tccattctgt gtttagatg ccagctgcat 120

gagaaaagg actcaccttc tggttcctg cctggaagag gaagagctgg cattgcacag 180

gagacggctg gacatgtctg aggcactgcc ctgccgggc aaggagacc ccacccagg 240

ctgcaggctg ggggcctgt attggcctg tgtccacaat gatccacc agtccaagc 300

catactggat ggtgggtct cccagagga gccacccag gtggacagca atgggagggt 360

gagatgtcct ggctcccag aacagctggg ggcattttg catccccc acccgctcct 420

ggcctggctc cctgagaggg gttcaggggc aatacctcct gc 462

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<210> 41

<211> 308

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 89 to 218

<400> 41

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ctctgggaca gatatgggtt tagagggtgc aaggggcct ggagtggccc agggggaaag 60

caggggatct gagctgccc tccctcagac aggcctcatg gtcgcatgct accacggctt 120

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ccagagtgtt gtggccctgc tcagccactg tcctttcctt gatgtgaacc agcaggacaa 180

aggaggggac acggccctca tgttggtgc ccaagcaggt gtgaggctgc tgcacccac 240

ttccgacagc ccccttttga tgcagacagg gcctcagccc cacccttggt gcacggtgtt 300

ctacacca 308

<210> 42

<211> 231

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 49 to 159

<400> 42

tcataccccc ctttcctggg gaccaagctt acccttgctg ccctgcaggc cacgtgcctc 60

tagtgagtct cctgctcaac tactatgtgg gcctggacct ggaacgccgg gaccagcggg 120

ggctcacggc gttaatgaag gctgccatgc ggaaccgctg tgagtgcgtg gccaccctcc 180

tcatggcagg tgtgcggggc ctggaccggg gtgtgtggcc tccagtcct c 231

<210> 43

<211> 231

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 49 to 189

<400> 43

tcataccccc ctttcctggg gaccaagctt acccttgctg ccctgcaggc cacgtgcctc 60

tagtgagtct cctgctcaac tactatgtgg gcctggacct ggaacgccgg gaccagcggg 120

ggctcacggc gttaatgaag gctgccatgc ggaaccgctg tgagtgcgtg gccaccctcc 180

tcatggcagg tgtgcggggc ctggaccggg gtgtgtggcc tccagtcctc c 231

<210> 44

<211> 588

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 98 to 499

<400> 44

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tatcccgccc catgtcaccc cctgtgctcc ttcccagggtg ctgacctgac agcagtggac 120

cctgttcggg gcaagacggc cctggaatgg gcagtgtga ccgacagctt cgacaccgtg 180

tggaggattc ggcagctgct gaggcggccc caagtggagc agcttagcca gcaactacaag 240

cccgagtggc cggccttgtc cgggctcgtg gcccaggccc aggccaggc ccaggttgcc 300

ccttcactcc tagaacggct gcaggctacc ttgagcctcc cctttgcccc gtctcctcag 360

gaggggggtg ttctggacca ctttgtgact gccacaacca gcttgccag tcccttcgtc 420

accactgcct gccacactct gtgccctgac catccacett cgctgggcac ccgaagcaag 480

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gggagtcccc agaggtcccc gtgggtcttc gtccctacc agagccct 588

<210> 45

<211> 503

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 27 to 503

<400> 45

ccaaggcatc ctcatcctcc caccagtgcc agccgaagcc cagtccttca ggacaccaaa 60

gtctggccct tcctctctgg cgataccagg agctcaggat agagaagagg aaacaggagg 120

aggaggccag aatggcacag aagtagggga agatgggata ggacaggctg ggaacaggta 180

atcaggcccc tcccaggget tctttccct ctggagtgcc tccggcctcc ccatccacct 240

ctgcctaagt aaatctgtc tcaacctata tatatacaag gtcattcatt ctagcattgt 300

ttgcaagagt gaaagagtgg aaacacccga agtgtccatc agtaagggaac aggctagatt 360

gattacggat gtaattgtg tccatccata cagagcatac tctacagtgt attctaaaat 420

aagactaagg aagctgttta tattctgata tgaaactacc atcaagatgt ataaagtaaa 480

aataactaag gagtggaaca gtg 503

25

49